# RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/585,923_

Source: _Two_

Date Processed by STIC: _2/2/07_

# ENTERED

IFWO

RAW SEQUENCE LISTING          DATE: 02/02/2007
PATENT APPLICATION:  US/10/585,923          TIME: 09:00:52

Input Set : A:\01-18-07 seq list_ST25.txt
Output Set: N:\CRF4\02022007\J585923.raw

```
 3 <110> APPLICANT: Brion, Aurelie
 4       Leissner, Philippe
 5       Berard, Cecile
 7 <120> TITLE OF INVENTION: Thrombosis Diagnosis/Prognosis Method
 9 <130> FILE REFERENCE: 128639
11 <140> CURRENT APPLICATION NUMBER: US 10/585,923
12 <141> CURRENT FILING DATE: 2006-07-13
14 <150> PRIOR APPLICATION NUMBER: PCT/FR05/50083
15 <151> PRIOR FILING DATE: 2005-02-10
17 <160> NUMBER OF SEQ ID NOS: 25
19 <170> SOFTWARE: PatentIn version 3.4
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 23
23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
26 <400> SEQUENCE: 1
27 aaattctcag aatttctgaa agg                                      23
30 <210> SEQ ID NO: 2
31 <211> LENGTH: 22
32 <212> TYPE: DNA
33 <213> ORGANISM: Homo sapiens
35 <400> SEQUENCE: 2
36 agtgcttaac aagaccatac ta                                       22
39 <210> SEQ ID NO: 3
40 <211> LENGTH: 22
41 <212> TYPE: DNA
42 <213> ORGANISM: Homo sapiens
44 <400> SEQUENCE: 3
45 agtgcttaac aagaccatac ta                                       22
48 <210> SEQ ID NO: 4
49 <211> LENGTH: 24
50 <212> TYPE: DNA
51 <213> ORGANISM: Homo sapiens
53 <400> SEQUENCE: 4
54 aacagacctg gaatttgaaa ctaa                                     24
57 <210> SEQ ID NO: 5
58 <211> LENGTH: 20
59 <212> TYPE: DNA
60 <213> ORGANISM: Homo sapiens
62 <400> SEQUENCE: 5
63 attactggct cttcctgagc                                          20
66 <210> SEQ ID NO: 6
67 <211> LENGTH: 20
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 68 <212> TYPE: DNA
 69 <213> ORGANISM: Homo sapiens
 71 <400> SEQUENCE: 6
 72 ttctgggctc ctggaaccaa                                             20
 75 <210> SEQ ID NO: 7
 76 <211> LENGTH: 20
 77 <212> TYPE: DNA
 78 <213> ORGANISM: Homo sapiens
 80 <400> SEQUENCE: 7
 81 tctagaaaca gttgcctggc                                             20
 84 <210> SEQ ID NO: 8
 85 <211> LENGTH: 20
 86 <212> TYPE: DNA
 87 <213> ORGANISM: Homo sapiens
 89 <400> SEQUENCE: 8
 90 ctaccagcgt gccaccaggt                                             20
 93 <210> SEQ ID NO: 9
 94 <211> LENGTH: 16
 95 <212> TYPE: DNA
 96 <213> ORGANISM: Homo sapiens
 99 <220> FEATURE:
100 <221> NAME/KEY: misc_feature
101 <222> LOCATION: (13)..(13)
102 <223> OTHER INFORMATION: n is inosine
104 <400> SEQUENCE: 9
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W--> 105 ctggacaggc gangaa                                               16
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108 <210> SEQ ID NO: 10
109 <211> LENGTH: 16
110 <212> TYPE: DNA
111 <213> ORGANISM: Homo sapiens
114 <220> FEATURE:
115 <221> NAME/KEY: misc_feature
116 <222> LOCATION: (13)..(13)
117 <223> OTHER INFORMATION: n is inosine
119 <400> SEQUENCE: 10
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W--> 120 ctggacaggc aangaa                                               16
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123 <210> SEQ ID NO: 11
124 <211> LENGTH: 18
125 <212> TYPE: DNA
126 <213> ORGANISM: Homo sapiens
128 <400> SEQUENCE: 11
129 actctcagca agcctcaa                                               18
132 <210> SEQ ID NO: 12
133 <211> LENGTH: 18
134 <212> TYPE: DNA
135 <213> ORGANISM: Homo sapiens
138 <220> FEATURE:
139 <221> NAME/KEY: misc_feature
140 <222> LOCATION: (13)..(13)
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     141  <223> OTHER INFORMATION: n is inosine
     143  <400> SEQUENCE: 12
W--> 144  actctcagcg agnctcaa                                        18
     147  <210> SEQ ID NO: 13
     148  <211> LENGTH: 51
     149  <212> TYPE: DNA
     150  <213> ORGANISM: Artificial Sequence
     152  <220> FEATURE:
     153  <223> OTHER INFORMATION: T7 Phage Polymerase Promoter
     155  <400> SEQUENCE: 13
     156  aattctaata cgactcacta tagggagaaa attctcagaa tttctgaaag g         51
     159  <210> SEQ ID NO: 14
     160  <211> LENGTH: 51
     161  <212> TYPE: DNA
     162  <213> ORGANISM: Artificial Sequence
     164  <220> FEATURE:
     165  <223> OTHER INFORMATION: T7 Phage Polymerase Promoter
     167  <400> SEQUENCE: 14
     168  aattctaata cgactcacta tagggagaag gattactggc tcttcctgag c         51
     171  <210> SEQ ID NO: 15
     172  <211> LENGTH: 20
     173  <212> TYPE: DNA
     174  <213> ORGANISM: Homo sapiens
     176  <400> SEQUENCE: 15
     177  agctgcccat gaatagcact                                        20
     180  <210> SEQ ID NO: 16
     181  <211> LENGTH: 21
     182  <212> TYPE: DNA
     183  <213> ORGANISM: Homo sapiens
     185  <400> SEQUENCE: 16
     186  ctggaaccaa tcccgtgaaa g                                      21
     189  <210> SEQ ID NO: 17
     190  <211> LENGTH: 18
     191  <212> TYPE: DNA
     192  <213> ORGANISM: Homo sapiens
     194  <400> SEQUENCE: 17
     195  actctcagca agcctcaa                                         18
     198  <210> SEQ ID NO: 18
     199  <211> LENGTH: 15
     200  <212> TYPE: DNA
     201  <213> ORGANISM: Homo sapiens
     203  <400> SEQUENCE: 18
     204  tctcagcggg cctca                                            15
     207  <210> SEQ ID NO: 19
     208  <211> LENGTH: 44
     209  <212> TYPE: DNA
     210  <213> ORGANISM: Artificial Sequence
     212  <220> FEATURE:
     213  <223> OTHER INFORMATION: T7 Phage Polymerase Promoter
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216 aattctaata cgactcacta taggagctgc ccatgaatag cact                44
219 <210> SEQ ID NO: 20
220 <211> LENGTH: 28
221 <212> TYPE: DNA
222 <213> ORGANISM: Homo sapiens
225 <220> FEATURE:
226 <221> NAME/KEY: misc_feature
227 <222> LOCATION: (19)..(19)
228 <223> OTHER INFORMATION: n is inosine
230 <400> SEQUENCE: 20
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W--> 231 cgatcgctgg acaggcgang aacgatcg                                 28

```
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235 <211> LENGTH: 28
236 <212> TYPE: DNA
237 <213> ORGANISM: Homo sapiens
240 <220> FEATURE:
241 <221> NAME/KEY: misc_feature
242 <222> LOCATION: (19)..(19)
243 <223> OTHER INFORMATION: n is inosine
245 <400> SEQUENCE: 21
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W--> 246 cgatcgctgg acaggcaang aacgatcg                                 28

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250 <211> LENGTH: 30
251 <212> TYPE: DNA
252 <213> ORGANISM: Homo sapiens
254 <400> SEQUENCE: 22
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258 <210> SEQ ID NO: 23
259 <211> LENGTH: 30
260 <212> TYPE: DNA
261 <213> ORGANISM: Homo sapiens
264 <220> FEATURE:
265 <221> NAME/KEY: misc_feature
266 <222> LOCATION: (19)..(19)
267 <223> OTHER INFORMATION: n is inosine
269 <400> SEQUENCE: 23
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W--> 270 cggtcgactc tcagcgagnc tcaacgaccg                                30

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274 <211> LENGTH: 30
275 <212> TYPE: DNA
276 <213> ORGANISM: Homo sapiens
278 <400> SEQUENCE: 24
279 cgatcgactc tcagcaagcc tcaacgatcg                                30
282 <210> SEQ ID NO: 25
283 <211> LENGTH: 27
284 <212> TYPE: DNA
285 <213> ORGANISM: Homo sapiens
287 <400> SEQUENCE: 25
```

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288 cgatcgtctc agcgggcctc acgatcg                                    27

**RAW SEQUENCE LISTING ERROR SUMMARY**
PATENT APPLICATION:  **US/10/585,923**

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**Please Note:**
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; N Pos. 13
Seq#:10; N Pos. 13
Seq#:12; N Pos. 13
Seq#:20; N Pos. 19
Seq#:21; N Pos. 19
Seq#:23; N Pos. 19

```
L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
```